

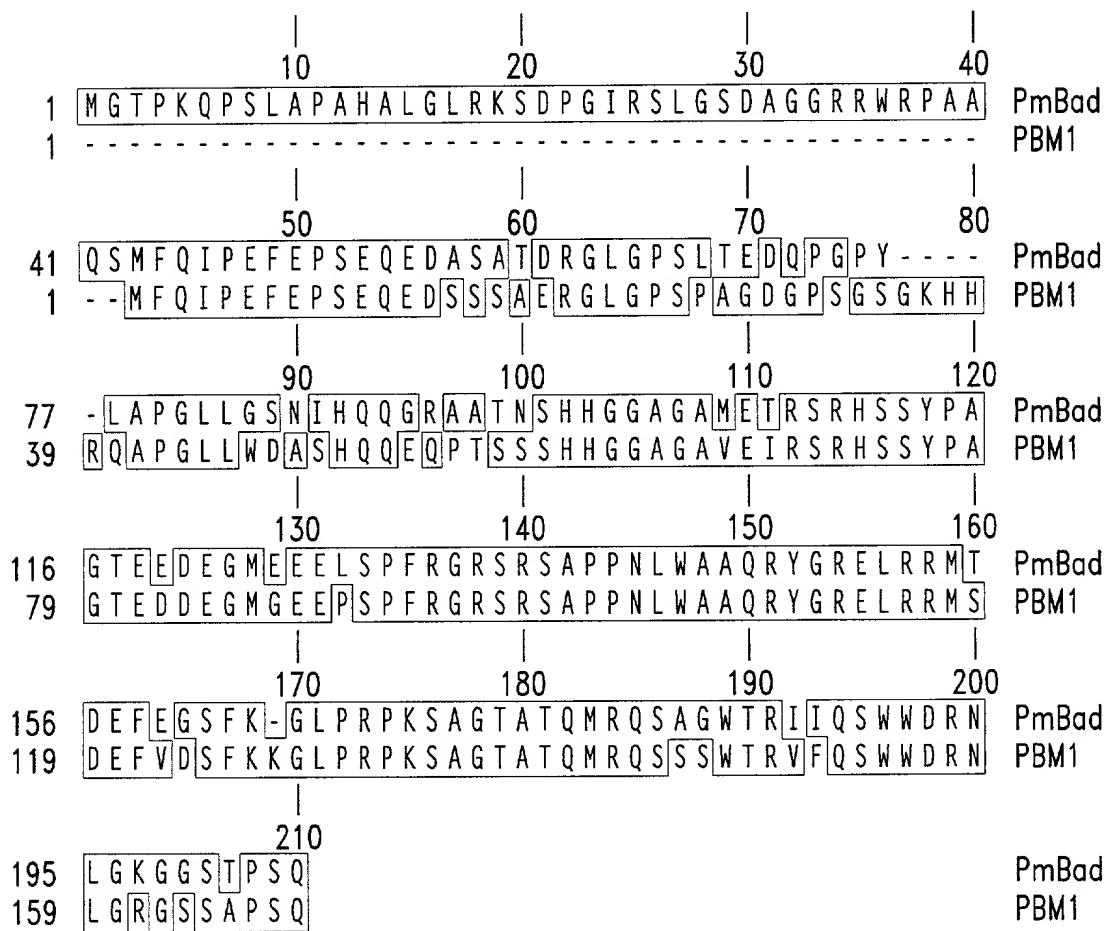
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Glu Arg Gly Leu Gly Pro Ser Pro Ala Gly Asp Gly Pro Ser Gly Ser	
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Gly Lys His His Arg Gln Ala Pro Gly Leu Leu Trp Asp Ala Ser His	
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Gln Gln Glu Gln Pro Thr Ser Ser Ser His His Gly Gly Ala Gly Ala	
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Val Glu Ile Arg Ser Arg His Ser Ser Tyr Pro Ala Gly Thr Glu Asp	
70 75 80	
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Asp Glu Gly Met Gly Glu Glu Pro Ser Pro Phe Arg Gly Arg Ser Arg	
85 90 95	
TCG GCG CCC CCC AAC CTC TGG GCA GCA CAG CGC TAT GGC CGC GAG CTC	393
Ser Ala Pro Pro Asn Leu Trp Ala Ala Gln Arg Tyr Gly Arg Glu Leu	
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Arg Pro Lys Ser Ala Gly Thr Ala Thr Gln Met Arg Gln Ser Ser Ser	
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Trp Thr Arg Val Phe Gln Ser Trp Trp Asp Arg Asn Leu Gly Arg Gly	
150 155 160	

Fig. 1A

Fig. 1B

[illegible]

ALIGNMENT OF MOUSE AND HUMAN BAD PROTEIN SEQUENCES



PmBad = mouse protein sequence
 PBM1 = human protein sequence

Fig. 2

Bad 2-Hybrid Interactions

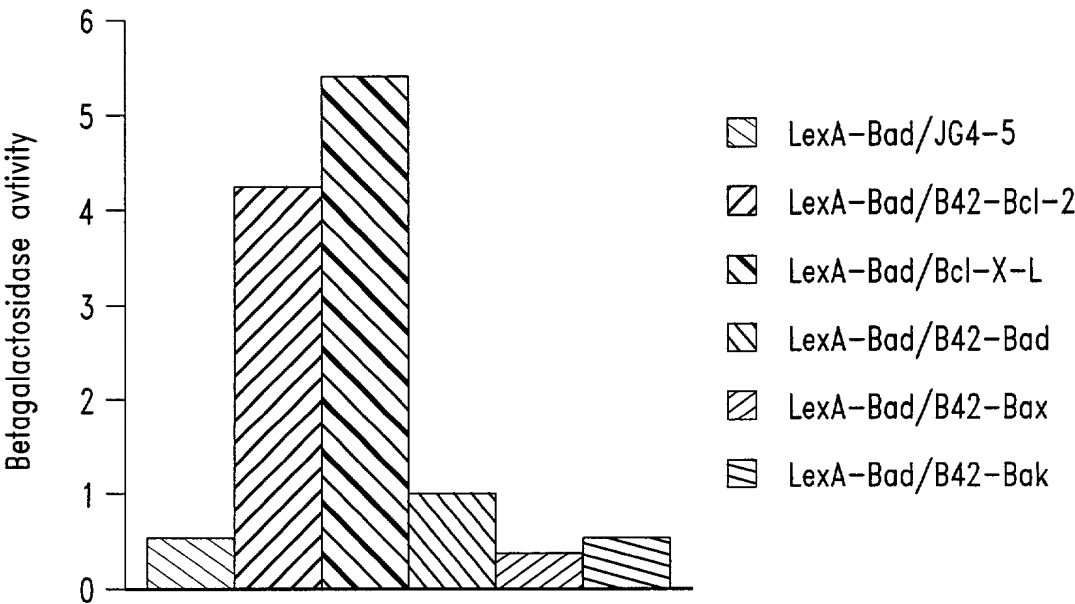


Fig. 3

Binding of Bcl-2 Family Proteins to Bad

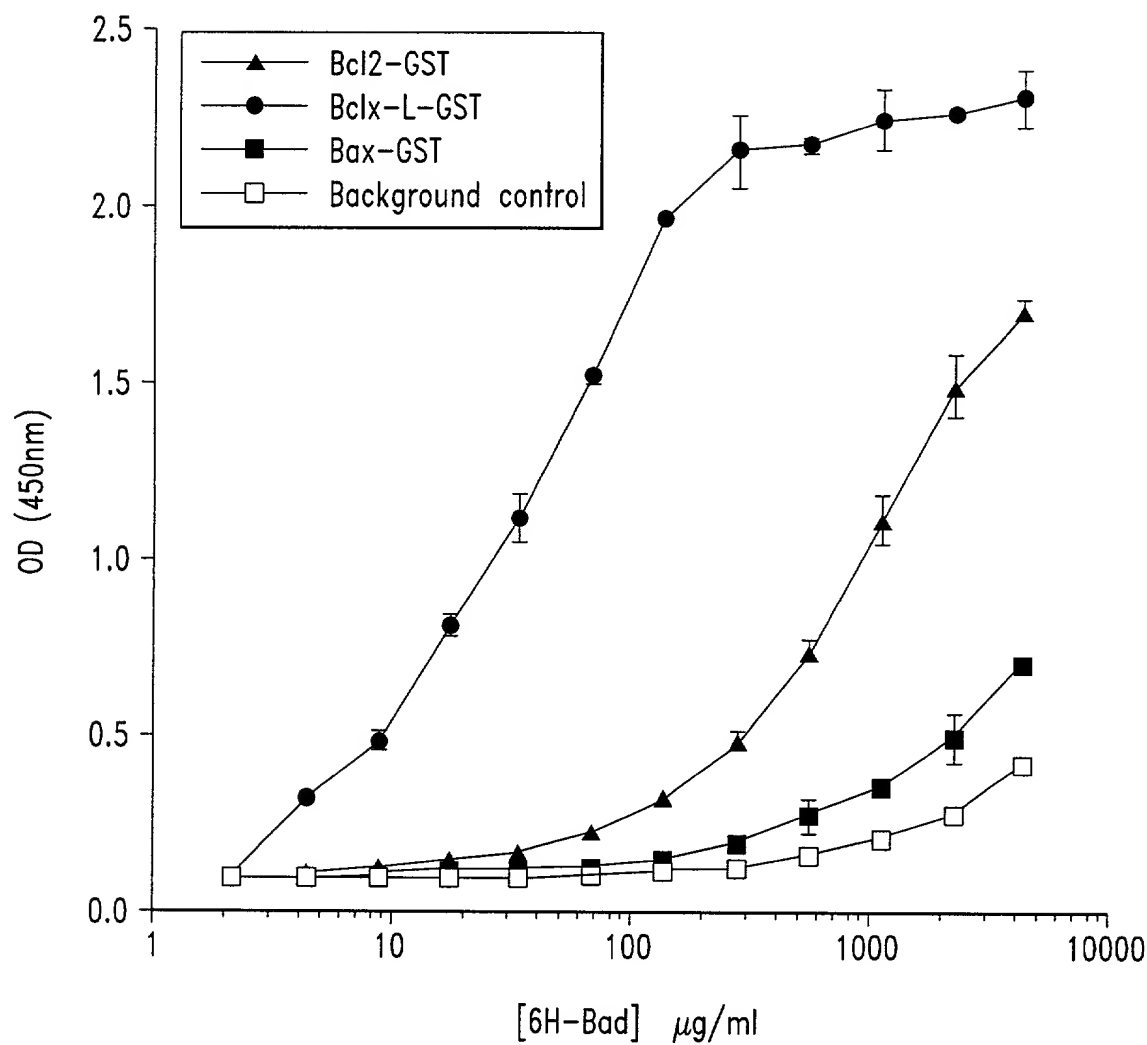


Fig. 4